### **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:  $\frac{10/568/396}{50 \text{ urce:}}$ Date Processed by STIC:  $\frac{2/27/06}{2}$ 

# ENTERED



IFWP

RAW SEQUENCE LISTING DATE: 02/27/2006
PATENT APPLICATION: US/10/568,396 TIME: 14:54:36

Input Set : A:\67489-PCT-US\_SeqList.txt
Output Set: N:\CRF4\02272006\J568396.raw

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4 <110> APPLICANT: Stephen P. Goff
             Guanxia Gao
      7 <120> TITLE OF INVENTION: ZAP PROTEIN AND RELATED COMPOSITIONS AND METHODS
      9 <130> FILE REFERENCE: 67489-PCT-US/JPW/JW
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/568,396
C--> 11 <141> CURRENT FILING DATE: 2006-02-13
     11 <150> PRIOR APPLICATION NUMBER: PCT/US2004/026162
     12 <151> PRIOR FILING DATE: 2004-08-12
     14 <160> NUMBER OF SEQ ID NOS: 10
     16 <170> SOFTWARE: PatentIn version 3.1
     18 <210> SEQ ID NO: 1
     19 <211> LENGTH: 776
     20 <212> TYPE: PRT
     21 <213> ORGANISM: mammalian
     23 <400> SEQUENCE: 1
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     34 Phe Val Leu Leu Glu Thr Gly Gly Gln Ala Gly Ile Thr Arg Ser Val
     37 Val Ala Thr Thr Arg Ala Arg Val Cys Arg Arg Lys Tyr Cys Gln Arg
                            70
     40 Pro Cys Asp Ser Leu His Leu Cys Lys Leu Asn Leu Leu Gly Arg Cys
                        85
     43 His Tyr Ala Gln Ser Gln Arg Asn Leu Cys Lys Tyr Ser His Asp Val
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                                        105
     46 Leu Ser Glu Gln Asn Phe Gln Ile Leu Lys Asn His Glu Leu Ser Gly
     47
                                    120
     49 Leu Asn Gln Glu Glu Leu Ala Cys Leu Leu Val Gln Ser Asp Pro Phe
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     52 Phe Leu Pro Glu Ile Cys Lys Ser Tyr Lys Gly Glu Gly Arg Lys Gln
                            150
                                                155
     55 Thr Cys Gly Gln Pro Gln Pro Cys Glu Arg Leu His Ile Cys Glu His
                                            170
     58 Phe Thr Arg Gly Asn Cys Ser Tyr Leu Asn Cys Leu Arg Ser His Asn
                                        185
                    180
     61 Leu Met Asp Arg Lys Val Leu Thr Ile Met Arg Glu His Gly Leu Ser
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     64 Pro Asp Val Val Gln Asn Ile Gln Asp Ile Cys Asn Asn Lys His Ala
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215

220

210

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67 Arg Arg Asn Pro Pro Gly Thr Arg Ala Ala His Pro His Arg Arg Gly 68 225 230 235 70 Gly Ala His Arg Asp Arg Ser Lys Ser Arg Asp Arg Phe Leu His Asn 250 73 Ser Leu Glu Phe Leu Ser Pro Val Val Ser Pro Leu Gly Ser Gly Pro 260 265 76 Pro Ser Pro Asp Val Thr Ser Cys Lys Asp Ser Leu Glu Asp Val Ser 275 280 79 Val Asp Val Thr Gln Lys Phe Lys Tyr Leu Gly Thr His Asp Arg Ala 290 295 300 82 Gln Leu Ser Pro Val Ser Ser Lys Ala Ala Gly Val Gln Gly Pro Ser 310 315 85 Gln Met Arg Ala Ser Gln Glu Phe Ser Glu Asp Gly Asn Leu Asp Asp 330 88 Ile Phe Ser Arg Asn Arg Ser Asp Ser Ser Ser Arg Ala Ser Ala 340 345 91 Ala Lys Val Ala Gln Arg Asn Glu Ala Val Ala Met Lys Met Gly Met 360 94 Glu Val Lys Gly Lys Lys Glu Ala Pro Asp Ile Asp Arg Val Pro Phe 375 380 97 Leu Asn Ser Tyr Ile Asp Gly Val Thr Met Glu Lys Ala Ser Val Ser 390 395 100 Gly Ile Pro Gly Lys Lys Phe Thr Ala Asn Asp Leu Glu Asn Leu Leu 405 410 103 Leu Leu Asn Asp Thr Trp Lys Asn Val Ala Lys Pro Gln Asp Leu Gln 420 425 106 Thr Thr Gly Arg Ile Thr Asp Ser Gly Gln Asp Lys Ala Phe Leu Gln 435 440 109 Asn Lys Tyr Gly Gly Asn Pro Val Trp Ala Ser Ala Ser Thr His Asn 455 112 Ala Pro Asn Gly Ser Ser Gln Ile Met Asp Glu Thr Pro Asn Val Ser 470 475 115 Lys Ser Ser Thr Ser Gly Phe Ala Ile Lys Pro Ala Ile Ala Gly Gly 485 490 118 Lys Glu Ala Val Tyr Ser Gly Val Gln Ser Pro Arg Ser Gln Val Leu 500 505 121 Ala Val Pro Gly Glu Ala Thr Thr Pro Val Gln Ser Asn Arg Leu Pro 520 515 124 Gln Ser Pro Leu Ser Ser Ser His Arg Ala Ala Ser Gly Ser 535 127 Pro Gly Lys Asn Ser Thr His Thr Ser Val Ser Pro Ala Ile Glu Ser 550 555 130 Ser Arg Met Thr Ser Asp Pro Asp Glu Tyr Leu Leu Arg Tyr Ile Leu 565 570 133 Asn Pro Leu Phe Arg Met Asp Asn His Gly Pro Lys Glu Ile Cys Gln 580 585 136 Asp His Leu Tyr Lys Gly Cys Gln Gln Ser His Cys Asp Arg Ser His 600 139 Phe His Leu Pro Tyr Arg Trp Gln Met Phe Val Tyr Thr Trp Arg RAW SEQUENCE LISTING DATE: 02/27/2006
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140
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142 Asp Phe Gln Asp Met Glu Ser Ile Glu Gln Ala Tyr Cys Asp Pro His
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145 Val Glu Leu Ile Leu Ile Glu Asn His Gln Ile Asn Phe Gln Lys Met
                                         650
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148 Thr Cys Asp Ser Tyr Pro Ile Arg Arg Leu Ser Thr Pro Ser Tyr Glu
                660
                                    665
151 Glu Lys Pro Leu Ser Ala Val Phe Ala Thr Lys Trp Ile Trp Tyr Trp
            675
                                680
                                                     685
154 Lys Asn Glu Phe Asn Glu Tyr Ile Gln Tyr Gly Asn Glu Ser Pro Gly
        690
                            695
157 His Thr Ser Ser Asp Ile Asn Ser Ala Tyr Leu Glu Ser Phe Phe Gln
158 705
                        710
                                             715
160 Ser Cys Pro Arg Gly Val Leu Pro Phe Gln Ala Gly Ser Gln Lys Tyr
161
                    725
                                         730
163 Glu Leu Ser Phe Gln Gly Met Ile Gln Thr Asn Ile Ala Ser Lys Thr
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166 Gln Arg His Val Val Arg Arg Pro Val Phe Val Ser Ser Asn Asp Val
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176 <213> ORGANISM: mammalian
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181 atgaccotgg aggaactgct gggtgagatc aggctccccg aggcgcagct ctacgagctg
                                                                          120
183 ctggagacgg cggggcccga tcgcttcgtg ctattggaga ctggaggcca ggccgggatc
                                                                          180
185 acteggtetg tagtggetac tactegagee egegtetgee gteggaagta etgeeagaga
                                                                          240
187 ccctgcgaca gcctgcacct ctgcaagctt aatctgctcg gccggtgcca ctatgcacag
                                                                          300
189 teteagegga acetetgeaa atatteteae gatgttetet eggaacagaa etteeagate
                                                                          360
191 ctgaagaatc atgagctctc tgggcttaac caagaggagc tagcttgcct cctggtccaa
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193 agcgaccctt ttttcctgcc cgagatatgc aagagttaca aaggagaggg ccgaaaacag
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195 acctgtgggc agccacagcc atgcgagaga ctccacatct gtgagcactt cacceggggc
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197 aactgcagtt acctcaactg tctcaggtct cacaacctga tggacagaaa ggtgttgacc
                                                                          600
199 atcatgaggg agcacgggct gagtcctgat gtggtccaga acatccagga catctgcaac
                                                                          660
                                                                          720
201 aacaaacacg ccaggaggaa cccgcctggc acgagagctg cccatccaca ccgcagaggc
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203 ggcgcacaca gagacagaag caaaagcaga gaccgcttcc ttcacaacag tctagaattt
205 ctctcacctg ttgtctcacc tctgggatct ggtccgccta gcccagatgt caccagctgt
                                                                          840
207 aaagattccc tggaggatgt gtctgtggat gtcacccaga agttcaagta cttggggacg
                                                                          900
209 catgaccgtg cgcagctctc cccagtctca tctaaggctg ctggtgttca aggacccagt
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211 caaatgagag caagccaaga gttttcagag gatgggaatc tagatgacat attttctagg
                                                                         1020
213 aatcgttctg attcatcatc aagtcgagcc tccgctgcca aggtggcaca aagaaatgaa
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215 gctgtggcca tgaaaatggg catggaggtc aagggcaaga aggaggctcc agacatcgat
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217 cgggtcccat ttttaaatag ttatattgat ggggtgacca tggaaaaagc atcggtctca
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219 ggaattccag gcaaaaagtt cacagccaat gatctggaaa atttgctatt acttaacgac
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221 acttggaaga atgtggctaa gccccaggat ctgcagacca caggcagaat cactgacagt
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223 ggccaagaca aggcattcct gcagaataaa tatggaggaa acccagtgtg ggcaagtgca
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	aaaagtagta ccagtggttt tgccataaaa cca				1500	
	tattctggag ttcagagtcc gagaagccag gto				1560	
	cctgtacaga gcaacaggct gcctcagtcg cc	_	_		1620	
	gcctctggga gccctggcaa gaactccacc ca				1680	
235	tcaaggatga catcagaccc cgatgagtat cto	cctacgct	acatcctaaa	tcctttattt	1740	
237	aggatggata atcatggccc gaaggaaatc tg	tcaggacc	atctgtacaa	gggctgtcaa	1800	
239	cagagecact gegacaggag teacttecat etg	gccctacc	ggtggcagat	gttcgtatat	1860	
241	accacttgga gggacttcca ggacatggag to	tatcgaac	aggcctattg	tgatccccac	1920	
243	gttgaactca ttttgataga aaaccatcag at	caatttcc	agaaaatgac	ctgtgactcc	1980	
	taccccatcc gacgcctctc cactccctca ta				2040	
247	gccaccaagt ggatttggta ttggaagaat ga	atttaatg	aatatatcca	gtatgggaat	2100	
249	gagageceag gecaeaceag etetgaeate aad	ctctgcgt	acctggagtc	tttcttccag	2160	
251	tcttgtccca ggggagtttt gccattccag gc	tggttcac	agaagtacga	gttaagcttc	2220	
253	caagggatga ttcagacaaa tatagcttcc aa	gactcaaa	ggcatgttgt	cagaaggcca	2280	
255	gtatttgttt cttcgaacga tgtggagcag aag	gagaagag	gtccagagtg	a	2331	
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278	<223> OTHER INFORMATION: PCR Primer					
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### RAW SEQUENCE LISTING

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Input Set : A:\67489-PCT-US\_SeqList.txt Output Set: N:\CRF4\02272006\J568396.raw

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322	<212> TYPE: DNA	
323	<213> ORGANISM: Artificial Sequence	
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326	<223> OTHER INFORMATION: PCR Primer	
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#### VERIFICATION SUMMARY

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DATE: 02/27/2006 TIME: 14:54:37

PATENT APPLICATION: US/10/568,396 TIME: 1

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Output Set: N:\CRF4\02272006\J568396.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date